

GenCore version 5.1.4_p5_4578
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OM nucleic - nucleic search, using sw model

Run on: June 2, 2003, 00:51:28 ; Search time 4674.33 Seconds
(without alignments)
16997.243 Million cell updates/sec

Title: US-09-596-194-59

Perfect score: 2730

Sequence: 1 gtcgaccacgcgtccgtcc.....aaaaaaaaaggggcggccgc 2730

Scoring table: OLIGO_NUC

Gapop 60.0 , Gapext 60.0

Searched: 2054640 seqs, 14551402878 residues

Word size : 400

Total number of hits satisfying chosen parameters: 9

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Listing first 45 summaries

Database : GenEmbl:
1: gb_ba:
2: gb_htg:
3: gb_in:
4: gb_om:
5: gb_ov:
6: gb_pat:
7: gb_ph:
8: gb_pl:
9: gb_pr:
10: gb_ro:
11: gb_sts:
12: gb_sy:
13: gb_un:
14: gb_vi:
15: em_ba:
16: em_fun:
17: em_hum:
18: em_in:
19: em_mu:
20: em_om:
21: em_or:
22: em_ov:
23: em_pat:
24: em_ph:
25: em_pl:
26: em_ro:
27: em_sts:

Search repeated in

GenSeq

US Pats

US PGPUBS

EST

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OM nucleic - nucleic search, using sw model

Run on: June 2, 2003, 00:51:28 ; Search time 3446.67 Seconds
(without alignments)
16997.243 Million cell updates/sec

Title: US-09-596-194-60

Perfect score: 2013

Sequence: 1 atggcccagctgttcctgcc.....ccctgcagctctgggtcacc 2013

Scoring table: OLIGO_NUC

Gapop 60.0 , Gapext 60.0

Searched: 2054640 seqs, 14551402878 residues

Word size : 400

Total number of hits satisfying chosen parameters: 9

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Listing first 45 summaries

Database : GenEmbl:
1: gb_ba:
2: gb_htg:
3: gb_in:
4: gb_om:
5: gb_ov:
6: gb_pat:
7: gb_ph:
8: gb_pl:
9: gb_pr:
10: gb_ro:
11: gb_sts:
12: gb_sy:
13: gb_un:
14: gb_vi:
15: em_ba:
16: em_fun:
17: em_hum:
18: em_in:
19: em_mu:
20: em_om:
21: em_or:
22: em_ov:
23: em_pat:
24: em_ph:
25: em_pl:
26: em_ro:
27: em_sts:

Search repeated in
GenSeq
US Pats
US PGPubs
EST